1/**%**33 FIG 1

U3 R U5	U3 R LTR
ENV	ENV
U3 R U5 GAG POL ENV LTR	POL ENV LTR
US GAG	RNA (VIRION) U5 GAG
U3 R U5	GENOMIC R

PROVIRAL DNA

2 / ¾ 33 FIG 2

10	
10 20 30 40 50 1234567890 1234567890 1234567890 1234567890	
GCTIATAGAA GGACCCCIAG TATGGGGIAA TCCCCICIGG GAAACCAAGC A Y R R T P S M G . S P L G N Q A L I E G P L V W G N P L W E T K P L . K D P . Y G V I P S G K P S	50
CCCAGIACIC AGCAGGAAAA ATAGAATAGG AAACCICACA AGGACATACT PVLSRKNRIG NLTRTYF QYSAGKIE.ETSQGHT PSTQQEK.NRKPHKDIL	100
TTCCTCCCCT CCAGATGGCT AGCCACTGAG GAAGGAAAAA TACTTTCACC PPLQMASH.GRKNTFT FLPSRWLATEEGKILSP SSPPDG.PLRKEKYFHL	150
TGCAGCIAAC CAACAGAAAT TACTIAAAAC CCITCACCAA ACCITCCACT C S . P T E I T . N P S P N L P L A A N Q Q K L L K T L H Q T F H L Q L T N R N Y L K P F T K P S T	200
TAGGCATIGA TAGCACCCAT CAGATGGCCA AATTATTATT TACTGGACCA R H H P S D G Q I I I Y W T R G I D S T H Q M A K L L F T G P . A L I A P I R W P N Y Y L L D Q	250
PFQNYQEDSQGCGGAAGGGCCA PFQNYQEDSQGL.SVP GLFKTIKKIVRGCEVCQ AFSKLSRR.SGAVKCAK	300
AAGAAATAAT K K . R N N E I	310

3/3433 FIG 3A

10 20 30 40 50	
<u>1234567890 1234567890 1234567890 1234567890 1234567890</u>	
CCCIGIAICI TIAACCICCI TGITAAGITT GICICTICCA GAATCAAAAC	50
PCIFNLL VKF VSSR IKT	
PVS LTSL LSL S L P E S K L	
LYL. PPC. VCLFQ NQN	
TGIAAAACIA CAAATTGITC TTCAAATGGA GCACCAGATG GAGTCCATGA	100
V K L Q I V L Q M E H Q M E S M T	100
. NY KLF FKWS TRW SP.	
CKTT NCS SNG APDG VHD	
CRIINCS SNG APDG VAD	
CTAAGATCCA CCGTGGACCC CTGGACCGC CTGCTAGCCC ATGCTCCGAT	150
KIH RGP L D R P A S P C S D	
LRST VDP WTG LLAH APM	
. DPPWTPGPAC.PMLRC	
. Drrwii Grac. i mbke	
GITAATGACA TIGAAGGCAC COCICCOGAG GAAATCICAA CIGCACAACC	200
V N D I E G T P P E E I S T A Q P	
LMT LKAP LPR KSQ LHNP	
H . R H P S R G N L N C T T	
CCTACTATEC CCCAATTCAG CGGGAAGCAG TTAGAGCGGT CATCAGCCAA	250
LLCPNSAGSS.SGHQPT	
Y Y A P I Q R E A V R A V I S Q	
PTMP QFS GKQ LERS SAN	
CCTCCCCAAC AGCACTIGGG TITTCCTGTT GAGAGGGGGG ACTGAGAGAC	300
SPT ALG FSC. EGG LRD	
PPQQHLGFPVERGD.ET	
LPNSTWVFLLRGGTERQ	
AGGACTACCT GGATTTOCTA GGCCAACGAA GAATCCCTAA GCCTAGCTGG	350
RTSWIS. ANE ESLS LAG	
G L A G F P R P T K N P . A . L G	
D. L DFL GQRR IPK PSW	

4 / 34 33 FIG 38

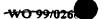
10		
10 20 1234567890 1234567890 1234567	30 40 900 1234567000 123456700	50
GAAGGIGACT GCATCCACCT CTAAACA K V T A S T S K H R . L H P P L N M E G D C I H L . T V	IGG GGCTTGCAAC TTAGCTCA(G A C N L A H G L A T L T	T 400
CCCGACCAAT CAGAGAGCIC ACTAAAAI RPIREL TKM PDQSESSLKC PTNQRAH N	L I R Q K . E	
GIAAAGAAAT AGCCAATCAT CIATIGOC V K K . P I I Y C L . R N S Q S S I A K E I A N H L L P	RAQREGQ.	
AGGATOGOGA TATAAACOCA GOCATTOG G S G Y K P R H S S D R D I N P G I R R I G I . T Q A F E	S R Q R Q P P I A G N G N P L	r 550
TGGGTCCCCT CCCTTGTAT GGGCGCTCT G P L P L Y G R S W V P S L C M G A L G S P P F V W A L C	V F T L F H S F S L Y F T L	
ATTANATOTT GCAACTGAAA AAAAAAAAAAAAAAAAAAAAAAAAAA	K K	635

5 / 34 33 FIG 4A

10 2 1234567890 123456789	0 30		50
ATGGCCCICC CTTATCATA M A L P Y H T W P S L I I G P P L S Y	C TTTTCTCTTT F L F L F S L	ACIGITCICT TACCCCC T V L L P P L F S Y P I	TTT 50 F
COCICICACT GCACCCCCI A L T A P P L S L H P L R S H C T P S	P C C C H A A V	TTSSSF QPVAP	Y L
ACCAAGAGIT TCTATGAAG Q E F L . R T K S F Y E E P R V S M K	T R L P R G F	G N I D A L E I L M P	P H
TCATATAGGA GITTATCIA S Y R S L S K H I G V Y L I . E F I .	G N S R E T P	T F T A H T P S L P T P	H
TATGCCCCC AACIGCIAT M P R N C Y C P A T A I Y A P Q L L .	N S A T T L P L	L C M H A N F A C M Q	T I
CICATIATIG GACAGGGAA H Y W T G K L I I G Q G K S L L D R E	M I N P	S C P G G L V V L E D	L L
GAGCACIG TCIGITGGA G A T V C W T E P L S V G S H C L L D	Y F T L T S P	H T S M S D	G f G

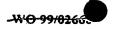
FIG 4B

10	20			
			1234567890 1234567890	
GGGTGGAATT	CAAGGTCAGG	CAAGAGAAAA	ACAAGTAAAG GAAGCAATCT	400
G G I	Q G Q A	R E K	Q V K E A I S	100
V E F	K V R	QEKN	K * R K Q S	
G W N S	R S G	KRK	T S K G S N L	
CCCAACTGAC	CCGGGGACAT	AGCACCCCTA	GCCCCTACAA AGGACTAGTT	450
O I T	R G H	S T P S	P Y K G L V	450
			APTKD*F	
	Б С ш +	ир +	P L Q R T S S	
F I D	r G i "	пр	PLQRTSS	
$C\Psi C\Psi C X X X X C$	TACATCAAA	CCTCCCTTCC	CITICAL ALLES	
			CATACTCGCC TGGTGAGCCT	500
			H T R L V S L	
5 Q N	Y M K P	S V P	I L A W * A Y	
L K T	T * N	PPYP	Y S P G E P	
ATTTAATACC	ACCCTCACTC	GGCTCCATGA	GGTCTCAGCC CAAAACCCTA	550
F N T	TLTR	L H E	V S A Q N P T	
L I P	P S L	G S M R	SQPKTL	
I * Y H	P H S	A P *	G L S P K P Y	
			TCAGGCCATA CATTTCAATC	600
$N \subset W$	M C L	P L H F	RPYISI	
L T V G	C A S	P C T	S G H T F Q S	
* L L	D V P P	P A L	Q A I H F N P	
CCTGTTCCTG	AACAATGGAA	CAACTTCAGC	ACAGAAATAA ACACCACTTC	650
			T E I N T T S	
			Q K * T P L P	
C S *	T M E	огон	RNKHHF	
-	- 	- · ·		
CGTTTTAGTA	GGACCTCTTG	TTTCCAATCT	GGAAATAACC CATACCTCAA	700
		SNL		, 00
			K.* P I P O	
RFSP	T S C	FOS	G N, N P Y L K	
	± 5 ¢	. <u>v</u> 3	ои,ие і ь К	



7/33 6/32 FIG 4 (continued) FIG 4C

)	10 20 30 40 50
)	1234567890 1234567890 1234567890 1234567890 1234567890
	ACCICACCIG TGIAAAATIT AGCAATACIA TAGACACAAC CAGCICCCAA L T C V K F S N T I D T T S S Q T S P V . N L A I L . T Q P A P N P H L C K I . Q Y Y R H N Q L P M
	TGCATCAGGT GGGTAACACC TCCCACAGGA ATAGICIGCC TACCCTCAGG C I R W V T P P T R I V C L P S G A S G G . H L P H E . S A Y P Q E H Q V G N T S H T N S L P T L R
850	AATATTTTT GICIGIGGIA CCICAGCCIA TCATTGITTG AATGCCICIT I F F V C G T S A Y H C L N G S S Y F L S V V P Q P I I V . M A L N I F C L W Y L S L S L F E W L F
900	CACAATCIAT GIGCITOCIC TCATICITAG TGCCCCCTAT GACCATCIAC E S M C F L S F L V P P M T I Y Q N L C A S S H S . C P L . P S T R I Y V L P L I L S A P Y D H L H
950	ACTGAACAAG ATTTATACAA TCATGTOGTA OCTAAGOOOC ACAACAAAAG T E Q D L Y N H V V P K P H N K R L N K I Y T I M S Y L S P T T K E . T R F I Q S C R T . A P Q Q K
1000	AGRACOCATT CITOCITTIG TRAICAGAGC AGGAGIGCIA GGCAGACIAG V P I L P F V I R A G V L G R L G Y P F F L L L S E Q E C . A D . S T H S S F C Y Q S R S A R Q T R
1050	GTACTOGCAT TOGCAGTATC ACAACCTCTA CTCAGTTCTA CTACAAACTA T G I G S I T T S T Q F Y Y K L V L A L A V S Q P L L S S T T N Y Y W H W Q Y H N L Y S V L L Q T I



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FIG 4 (continued)

					
			40		
<u>1234567890</u>	1234567890	1234567890	1234567890	1234567890	
S Q E I L K K	N G D . M V T	M E Q W N R	GICACIGACT V T D S S L T H . L	L V T P W S P	1100
L Q D C K I	Q L N S N L T	L A A P . Q Q	AGIAGICCIT V V L . S F S S P S	Q N R R K I E	1150
A L D T	L L T C . P	A K R G P K E	GGGGAACCIG ' G T C G E P V G N L 1	L F L Y F .	1200
G E E R E K N	C Y Y A V I M	V N Q L I N	TCCACAATIG S R I V P E L S Q N C	T E K S L R K	1250
V K E L K K	I R D R F E I	I Q C E Y N V	TAGAGCAGAG (R A E I E Q R . S R G	E L Q N S F K	1300
TER	W G L G A S	LSQW SAN	GGATGCCCTG (M P W G C P G D A L (V L P F S P	1350
TICTIAGGAC (F L G P S . D :	L A A L . Q L	L I L . Y C		G P C L D P V	1400

9|33 8|32 FIG 4 (continued) FIG 4 E

10 20 30	40 50	
1234567890 1234567890 1234567890		
TATCITIAAC CICCIIGITA AGITIGICIC II F N L L V K F V S S L T S L L S L S L Y L . P P C . V C L I	TICCAGAATT GAAGCIGIAA S R I E A V K P E L K L . F Q N . S C K	1450
AGCTACAGAT GGTCTTACAA ATGGAACCCC A LQMVLQMEP SYRWSYKWNP ATDGLTNGTP	A	1481

10133 -9/32 FIG 5 A

10 20				
1234567890 1234567890 1 TCAAAATCGA AGAGCTTTAG A				
S K S K S F R				50
QNRRALD				
KIE EL. 1				
			0 1 1	
GITTATTITT AGGGGAAGAA 1	CCIGITAGI	AIGITAATCA	ATCIGGAATC	100
F I F R G R M				
LFLGEE	C C . Y	V N Q	SGI	
V Y F . G K N	AVS	MLIN	L E S	
ATTACTGAGA AAGTTAAAGA A				150
Y . E S . R N				
ITEKVKE				
LLRKLKK	F E I	EIN	VEQR	
GGACCITCAA AACACIGCAC C	CIGGGGCCT (CCTCAGCCAA (IGGAIGCCT	200
G P S K H C T				
DLQNTAP		_ _		
TFK TLH P				
GEACTICICOC CITICITAGGA C				250
DSPLLRT				
TLPFLGP				
GLSPS.D	L . Q	L . Y F	YSS	·
TTTGCACCCT GIATCITCAA C	TROCHIGHT	AAGITIGICT (CTTCCAGAAT	300
WTLYLQL				
FGPCIFN			-	
LDPVSST	S L L	S L S	L P E L	
				4
TGAAGCIGIA AAGCIACAAA T				350
SCKATN				
EAVKLQI				
KL.SYK.	r r K	W IN P	r C S	

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FIG 5 (continued)

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
			ACCGGCCTGC P A C		400
мтк	ΙΥR	G P L D	R P A	R L C	
			T G L L		
TCIGATGITA	ATGACATIGA	AGTCACCCCT	CCCCAGGAAA	TCTCAACTGC	450
. c .	. н .	S H P S	R G N	LNC	
			P E E I		
·L M L	M T L K	S P L	P R K	S Q L H	
ACAACCCTA	CTACACTCCA	ATTCAGTAGG	AAGCAGTTAG	AGCAGITIGIC	500
• •			K Q L E		
			SS.		
			A V R		
عصعمصت	CCAACACTA	CTTGGGTTTT	CCIGITGAGA	GCCTCCACTG	550
A N L	PNST	WVF	L L R	GWTE	
			C . E		
			P V E R		
7C7C7C7C7	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	شبيكيس كيس	GACTAAGAAT	CCNDAGCTT	600
			L R I		
R D R T	. L W T	S A	D . E S	XSL	
ETG	LAGF	PRL	TKN	PXAX	
ANCTOGGAAG					650
			TWGL		
X G K					
L G R	. P H	PSLN	M G L	A T .	
			AAATGCTAAT		700
			N A N		
H T R	PIR	E L T K	MLI	RQK	
L T P D	QSE	S S L	K C . S	G K N	



12/33 11/32 FIG 5 (continued) FIG 5C

10 20 30 40 50	
<u>1234567890 1234567890 1234567890 1234567890 1234567890</u>	
CAGGAGGIAA AGCAATAGOC AATCATCIAT TGCCTGAGAG CACAGCGGGA G G K A I A N H L L P E S T A G Q E V K Q . P I I Y C L R A Q R E R R . S N S Q S S I A . E H S G K	750
AGGACAAGGA TIGGGATATA AACTCAGGCA TICAAGCCAG CAACAGCAAC R T R I G I . T Q A F K P A T A T G Q G L G Y K L R H S S Q Q Q P D K D W D I N S G I Q A S N S N	800
P F G S P P I V W E L C F H S I S P L G P L P L Y G S S V F T L F P L W V P S H C M G A L F S L Y F	850
CACTCTATTA AATCATGCAA CIGCACTCTT CIGGICCGIG TTTTTTATGG LY. IMQLHSSGPCFLW HSIKSCNCTLLVRVFYG TLLNHATALFWSVFFMA	900
CTCAAGCTGA GCTTTTGTTC GCCATCCACC ACTGCTGTTT GCCACCGTCA L K L S F C S P S T T A V C H R H S S . A F V R H P P L L F A T V T Q A E L L F A I H H C C L P P S	950
CAGACCOGCT GCTGACTTCC ATCCCTTTGG ATCCAGCAGA GTGTCCACTG R P A A D F H P F G S S R V S T V D P L L T S I P L D P A E C P L Q T R C . L P S L W I Q Q S V H C	1000
TECTOCTEAT COAGCEAGGT ACCOCATTGOC ACTOCOGATO AGGCTAAAGG LLIQRGTHCHSRSG.R CS.SSEVPIATPDQAKG APDPARYPLPLPIRLKA	1050



13|33 12 / 32 FIG 5 (continued) FIG 5D

10 20 30 40 50 1234567890 1234567890 1234567890 1234567890	
CHGCATG TICCIGCATG GCTAAGIGCC TGGGTTIGIC CTAATAGAAC LAIVPAW LSAWVCPNRT LPLFLHG.VPGFVLIEL CHCSCMAKCLGLSN	1100
TGAACACTGG TCACTGGGTT CCATGGTTCT CTTCCATGAC CCACGGCTTC E H W S L G S M V L F H D P R L L N T G H W V P W F S S M T H G F . T L V T G F H G S L P . P T A S	1150
TAATAGAGCT ATAACACTCA CCGCATGGCC CAAGATTCCA TTCCTTGGTA I E L . H S P H G P R F H S L V S Y N T H R M A Q D S I P W Y N R A I T L T A W P K I P F L G I	1200
TCIGICAGGC CAAGAACCCC AGGICAGAGA ANGIGAGGCT TGCCACCATT S V R P R T P G Q R X . G L P P F L . G Q E P Q V R E X E A C H H L C E A K N P R S E X V R L A T I	1250
TGGGAAGIGG CCCACTGCCA TTTTGGTAGC GGCCCACCAC CATCTTGGGA G K W P T A I L V A A H H H L G S G S G P L P F W . R P T T I L G W E V A H C H F G S G P P P S W E	1300
CCIGIGGGAG CAAGGAICCC CCAGIAACA CGSKDPPVT AVGARIPQ. LWEQGSPSN	1329

|4||33 |13||32| |FIG 6A

10 20 30 40 50 1234567890 1234567890 1234567890 1234567890 1234567890 CCTAGAACGT ATTCTGGAGA ATTGGGACCA ATGTGACACT CAGACGCTAA 50 PRTYSGELGPM. HSDAK LERILEN WDQ CDT QTLR . NV FWR IGTN VTL RR. GAAAGAAAG ATTTATATIC TICIGCAGIA CCGCCIGGCC ACAATATCCT 100 KET IYIL LQY RLA TISS KKR FIF FCST AWPQYP ERND LYS SAV PPGH NIL CITICAAGGGA GAGAAACCIG GCITICCIGAG GGAAGTATAA ATTATAACAT SRERNLAS.GKYKL.H L Q G R E T W L P E G S I N Y N I FKGEKPGFLREV. IITS CATCTIACAG CTAGACCICT TCTGIAGAAA GGAGGGCAAA TGGAGTGAAG 200 HLTARPLL.KGGQME.S I L Q · L D L F C R K E G K · W S E V SYS . TSSVER RANGVK TGCCATATGT GCAAACTTTC TTTTCATTAA GAGACAACTC ACAATTATGT 250 AIC ANFL FIK RQL TIM. PYV QTF FSLR DNS QLC CHMC KLS FH. ETTH NYV AAAAAGIGIG GITTIATGCCC TACAGGAAGC CCTCAGAGIC CACCICCCTA 300 KVW FMP YRKP SES TSL K K C G L C P T G S P Q S P P P Y KSVVYALQEALRVHLPT CCCAGGIC CCCCCCCCA CICCITCCIC AACTAATAAG GACCCCCTT 350 PQRP L P D S F L N . . G P P F PSV PSPT PSS TNK DPPL PAS PPR L L PQ L I R T P L TAACOCAAAC GGTOCAAAAG GAGATAGACA AAGGGGTAAA CAATGAACCA NPNGPKG DRQ RGKQ. TK TQT VQK EIDK GVN NEP . PKR SKR R. TKG. TMNQ AAGAGIGOCA ATATIOCOCC ATTATICCCC CICCAAGCAG IGAGAGGAG 450 ECQ YSP IMPP PSS ERR K S A N I P R L C P L Q A V R G G RVPIPPD YAP SKQ . E E E ACAATICGC COAGCOAGAG TOCCIGIACC TITTICICIC TOAGACTIAA 500 RIRP SQS ACT FFSL RLK EFG PARV PVP FSL SDLK NSAQPECLYLFLSQT.



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FIG 6 (continued)

10 20 30 40 50	
1234567890 1234567890 1234567890 1234567890 1234567890 AGCAAATTAA AATAGACCTA GGTAAATTCT CAGATAACCC TGACGCTAT A N N R P R I L R P R L Y	550
Q I K I D L G K F S D N P D G Y S K L K . T . V N S Q I T L T A I	
ATTCATCTT TACAAGGGT AGGACAATCC TTTCATCTCA CATGCAGAGA . C F T R V R T I L . S D M E R I D V L Q G L G Q S F D L T W R D L M F Y K G . D N P L I . H G E I	600
TATAATGITA CTACTAAATC AGACACTAAC CCCAAATGAG AGAAGICCCG Y N V T T K S D T N P K . E K C R I M L L L N Q T L T P N E R S A A . C Y Y . I R H . P Q M R E V P	650
CTGTAACTOC AGCCGAGAG TTTGGGATC TTTGGTATCT CAGTGAGGCC C N C S P R V W R S L V S Q S G Q V T A A R E F G D L W Y L S Q A L . L Q P E S L A I F G I S V R P	700
AACAATAGCA TGACAACAGA GGAAAGAACA ACTOOCACAG GOCAGCAGCC Q D D N R G K N N S H R P A G N N R M T T E E R T T P T G Q Q A T I G . Q Q R K E Q L P Q A S R Q	750
AGITOCCAGT GEAGACCIC ATTOCCACAC AGAATCAGAA CATOCAGATT S S Q C R P S L G H R I R T W R L V P S V D P H W D T E S E H G D W F P V . T L I G T Q N Q N M E I	800
GGTGCCACAA ACATTIGCTA ACTIGCGTGC TAGAAGGACT GAGGAAAACT V P Q T F A N L R A R R T E E N . C H K H L L T C V L E G L R K T G A T N I C . L A C . K D . G K L	850
AGGAGAGC CHATGAATTA CHCAATGATG TOCACHATAA CACAGGGAAA E E A Y E L L N D V H Y N T G K R K K P M N Y S M M S T I T Q G K G R S L . I T Q . C P L . H R E R	900
GCAACAAAAT CTTACTGCTT TTCTGCACAG ACTAAGGCAG GCATTGAGGA G R K S Y C F S G Q T K G G I E E E E N L T A F L D R L R E A L R K K K I L L L F W T D . G R H . G	950
AGCATACCTC CCIGICACCT CACTCTATTG AAGGCCAACT AATCTTAAAG A Y L P V T . L Y . R P T N L K G H T S L S P D S I E G Q L I L K S I P P C H L T L L K A N . S . R	1000



16/33 15 / 32 **FIG 6** (continued) FIG 6C

10 20 30 4 0 50 1234567890 1234567890 1234567890 1234567890	
CATAAGITTA TCACTCAGTC ACCTCCAGAC ATTAGAAAAA ACTTCAAAAG	
. V Y H S V S C R H . K K L Q K D K F I T Q S A A D I R K N F K S I S L S L S Q L Q T L E K T S K V	1050
TCTGCCTTAG GCCCGCAGCA GAACTTAGAA ACCCTATTTA ACTTGGCATC S A L G P E Q N L E T L F N L A S L P . A R S R T . K P Y L T W H P C L R P G A E L R N P I . L G I	1100
CICAGITITT TATAATAGAG ATCAGGAGGA GCAGGOGAAA COGGACAAAC S V F Y N R D Q E E Q A K R D K R Q F F I I E I R R S R R N G T N L S F L R S G G A G E T G Q T	1150
GOGATAAAAA AAAAAAGGGG GGTCCACTAC TITAGTCATG GCCCTCAGCC D K K K R G G P L L . S W P S G G I K K K G G V H Y F S H G P Q A G . K K K G G S T T L V M A L R Q	1200
AACCACACTT TOGAGOCTCT GCAAAAGGGA AAAGCTGGGC AAATCAAATG KQTLEALQKG KAGQIKC SRLWRLC KRE KLG KSNA ADF GGS AKGKSWA NQM	1250
CCTAATAGGG CTGGCTTGCA GTGGGGTCTA CAAGGACACT TTAAAAAACA LIG LASS AVY KDT LKKI G WLP VRST RTL . KR PNRA GFQ CGL QGHF KKD	1300
TEATOCAAGT AGAAATAAGC COCCCCTTG TOCATGCCCC TEAGGICAAG I Q V E I S R P L V H A P Y V K L S K . K . A A P L S M P L T S R Y P S R N K P P P C P C P L R Q G	1350
GENATURCIE CRAGOCCAC TOCCOCAGE CRICARCATA CICICAGICA G I T G R P T A P G D E D T L S Q E S L E G P L P Q G M K I L . V R N H W K A H C P R G . R Y S E S	1400
CAACCOATTA ACCACATCAT COACCACCAG CACTGAGGGT COOCCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	1450
AGGGCAGC CATGCCATCA COCTCACAGA GODDGGGTA TGITTGACCA R Q P M P S P S Q S P G Y V . P S A S P C H H P H R A P G M F D H A P A H A I T L T E P R V C L T I	1500

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FIG (continued)

10 12345 <i>6</i> 7890	20 1234567890	30 1234567890	40 1234567890	50 1234567890	19
TTGAGAGCCA L R A	A				1511
. E P					
E S Q					

18133 17/32 FIG 7A

10 20 30 40 50	
1234567890 1234567890 1234567890 1234567890 1234567890 ATGGGCACA GCCATCATCA TRANSPORT	<u> </u>
ATGGGCAGCA GCCATCATCA TCATCATCAC AGCAGGGGCC TGGTGCCGGG M G S S H H H H H H S S G L V P R	50
CCGCAGCCAT ATCGCTACCA TCACTGGTGG ACAGCAAATG GGTCGGATCC G S H M A S M T G G Q Q M G R I L	100
TACAACCIAT TCICCACAAT TCCCACCAAT GICACACICA CACCCIAACA E R I L E N W D Q C D T Q T L R	150
AAGAAACGAT TTATATTCTT CTGCAGTACC GCCTGGCCAC AATATCCTCT K K R F I F F C S T A W P Q Y P L	200
TCAAGGGAGA GAAACCTGGC TTCCTGAGGG AAGTATAAAT TATAACATCA Q G R E T W L P E G S I N Y N I I	250
TCTTACAGCT AGACCICTTC TGTAGAAAGG AGGGCAAATG GAGTGAAGTG LQLDLFCRKEGKW'S EV	300
CCATATGICC AAACTTICIT TICATTAAGA GACAACTCAC AATTATGIAA PYVQTFFSLRDNSQLCK	350
AAAGIGIGGI TIATGOOCIA CAGGAAGOOC TCAGAGICCA CCICCCIACC K C G L C P T G S P Q S P P P Y P	400
CCAGGGICCC CICCOCCACT CCTTCCICAA CTAATAAGGA CCCCCTTTA S V P S P T P S S T N K D P P L	450
ACCCAAACCG TOCAAAACGA GATAGACAAA GGGGTAAACA ATGAACCAAAA T Q T V Q K E I D K G V N N E P K	500
CAGIGOCAAT ATTOCCOCAT TATGCCCCT CCAAGCAGIG AGAGCAGGAG S A N I P R L C P L Q A V R G G E	550
AATTOGGOOC AGOCAGAGIG OCTGIACCTT TITICICICIC AGACTTAAAG F G P A R V P V P F S L S D L K	600
CAAATTAAAA TACACCIAGG TAAATTCICA GATAACCCIG ACGGCTATAT Q I K I D L G K F S D N P D G Y I	650
TCATGITTIA CAAGOGITAG CACAATOCIT TCATCICACA TOCACACATA D V L Q G L G Q S F D L T W R D I	700
TAATGITACT ACTAAATCAG ACACTAACCC CAAATGAGAG AAGIGCOCCT M L L N Q T L T P N E R S A A	750
GTAACIGCAG COCCAGAGIT TOGGGATCIT TOGGGATCICA GICAGGCCAA V T A A R E F G D L W Y L S Q A N	800



19|33 -18 / 32 FIG 7 (continued) FIG TB

10 20 30 40 50	
1234567890 1234567890 1234567890 1234567890 1234567890	
CAATAGGATG ACAACAGGG AAAGAACAAC TOOCACAGGC CAGCAGGCAG	
N R M T T E E R T T T E E R T T T T T T T T T	850
NRM TTEERTT PTG QQAV	
THIS ACTOR ACTOR	
TICCCAGIGI AGACCCICAT TOGGACACAG AATCAGAACA TOGAGATTOG	900
PSV DPH WDTE SEH GDW	
TGCCACAAAC ATTTGCTAAC TTGCGTGCTA GAAGGACTGA GGAAAACTAG	950
CHKHLLT CVL EGLR KTR	950
GAAGAAGCCT ATGAATTACT CAATGATGTC CACTATAACA CAGGGAAAGG	
K K P M N V C M M C T T T	1000
K K P M N Y S M M S T I T Q G K E	
AACAAAAMIN MAGAAAAAA	
AACAAAATCT TACTGCTTTT CTGCACAGAC TAAGGGAGGC ATTGAGGAAG	1050
ENL TAF LDRL REALRK	•
CATACCICCC TGICACCIGA CICTATIGAA GGCCAACTAA TCTTAAAGGA	1100
HTSL SPD SIE GQLI LKD	1100
TAAGITTATC ACTCAGTCAG CTGCAGACAT TAGAAAAAC TTCAAAAGTC	1150
KFI TQSA ADI RKN FKSL	1150
TECCTAACCT TECCECCCA CTCCACCACC ACCACCACCA CCACTGAGAT	
P K I A A A A TOTAL ACACCA CCACICAGAT	1200
PKLAAALEHH HHH H.D	
COGCUCUTA ACAAAGCCCG AAAGCAAGCT GAGTIGGCIN GIGGONA	1247
PAAN KAR KEA ELAX G	



20|33 19/32 FIG 8A

10 20 30 40 50 1234567890 1234567890 1234567890 1234567890	
ATGCTACCA TGACTGGTGG ACAGCAAATG GGTGGGATCC TAGAAGGTAT	50
	50
M A S M T G G Q Q M G R I L E R I	
TCTGCAGAAT TGGGACCAAT GTGACACTCA GACGCTAAGA AAGAAACGAT	100
LEN W D Q C D T Q T L R K K R F	
TTATATTCIT CIGCAGIACC GCCIGGCCAC AATATCCICI TCAAGGGAGA I F F C S T A W P Q Y P L Q G R	150
·	
GAAACCTGGC TICCTGAGGG AAGIATAAAT TATAACATCA TCTTACAGCT E T W L P E G S I N Y N I I L Q L	200
AGACCICTIC TGIAGAAAGG AGGGCAAATG GAGTGAAGTG CCATATGTGC D L F C R K E G K W S E V P Y V Q	250
AAACTITCIT TICATTAAGA GACAACTCAC AATTATGTAA AAAGTGTGGT	300
T F F S L R D N S Q L C · K K C G	
TTATGCCCIA CAGGAAGCCC TCAGAGTCCA CCTCCCTACC CCAGCGTCCC	350
L C P T G S P Q S P P P Y P S V P	
-	
CTCCCCCACT CCTTCCTCAA CTAATAAGGA CCCCCTTTA ACCCAAACGG	400
	400
S P T P S S T N K D P P L T Q T V	
TOCAAAAGGA GATAGACAAA GGGGTAAACA ATGAACCAAA GAGTGCCAAT	450
Q K E I D K G V N N E P K S A N	
Q N D I D N G V N N D P N S A N	
ATTICCCCCAT TATGCCCCCT CCAAGCAGIG AGAGGAGGAG AATTICGGCCC	500
I P R L C P L O A V R G G E F G P	
	CEO.
AGCCACAGIG CCIGIACCIT TITCICICIC ACACTIAAAG CAAATIAAAA	550
ARV PVPF SLS DLK QIKI	•
TAGACCTAGG TAAATTCTCA GATAACCCTG ACGCCTATAT TGATGTTTTA	600
D L G K F S D N P D G Y I D V L	000
D D G R F S D N P D G I I D V L	
CAMEGGITAG CACAATOCIT TCATCTCACA TGCACACATA TAATGITACT	650
QGLG QSF DLT WRDI MLL	
ACTION A STOCKED A CONCESSION OF A A MICHAEL A A CONCESSION OF THE ACTION OF	700
ACTAAATCAG ACACTAACCC CAAATGAGAG AAGTGCCGCT GTAACTGCAG	700
LNQTLTPNERSAAVTAA	
	•
CCCGAGAGIT TGGCGATCIT TGGTATCICA GICAGGCCAA CAATAGGATG	750 ´
REFGDL WYLS QAN NRM	-
101100000000000000000000000000000000000	
ACAACAGAGG AAAGAACAAC TOOCACAGGC CAGCAGGCAG TTOOCAGTGT	800
TTEERTT PTG QQAV PSV	

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FIG 8 (continued)

10	20	30	40	50		
				1234567890		
AGACCCICAT	TGGGACACAG	AATCAGAACA S E H	TGGAGATTGG	TGCCACAAAC	850	
		GAAGGACIGA E G L R			900	
		CACTATAACA T I T			950	
		TAAGGGAGGC R E A	_	CATACCTCCC H T S L	1000	
	CICIATIGAA S I E	GCCAACTAA G Q L I	TCTTAAAGGA L K D		1050	
		TAGAAAAAAC R K N			1100	
	CTOGAGCACC L E H H	ACCACCACCA (H H H		CCGGCTGCTA P A A N	1150	
	AAAGGAAGCT K E A	GAGITIGOCIG (E L A G	GIGGCA G		1186	

22/33 21 / 32 FIG 9A

		· · · · · · · · · · · · · · · · · · ·		
	20			
	CCTGATC CAGCA	CAGGC GCCCATTC G A H C Q A P I A	CC TCTCCCAATT L S Q L A S P N W	50
	AIV PA PL FL	AQLSA HS.V	WVHP PGFI	
CTAATCGAGC TGA N R A E L I E L N	H.SL TSHW	G S T V V P R F	L F H D S S M T	150
		S L H G T H C M V	P R F H Q D S I	200
	V R P R T	P G Q R P Q V R	T Q G L E H K A	250
TGCCACCATG TTG PPCW CHHVG	K Q P T S S P P	TILE.	A A R H Q P A T	300
		PR.Q	F G D H L V T T	350
CGAAGGCACC TGA E G T K G P E R R D L N	IRNHE SATM	G I S K K G S P :	A I G N K Q L E	400



23 33

FIG 9 B

			40		
M F L P	CAAGGCAAAA K A K R Q K	ATGCCCCTAA M P L R C P .	C I L	GCACAATIOG E N W R I G	450
GACCAATTTG A D Q F D T N L S P I .	P Q T T L R Q	V R K . E K		F F C Y S S A	500
	L A T I W P R	S S S Y P L Q	R G R G G E	N L A S T W P	550
PEGS	Y K L I N Y	. H H L N T I	T A R	P V L L F C	600
R K G	QME GKWS	. S A E V P	ATATTTACAA IFTN YLQ IYK	F L F T F F S	650
	Q L A I N S Q	M L T L C . Q	V . F C D L	V F L H C S Y	700
TEAL	Q I L R F Y	L P T P S P P	G I S P A S P	PES	750
CICCCCAACT L P N L S P T P Q L	Y				764

24 | 33 23 | 32 FIG 10 A

			40 1234567890 12345678	
TGICCOCTGT C P L C V R C	GCTCCTGATC S . S A P D P	CAGCACAGGC S T G A Q A	GCCCATIGCC TCTCCCAA A H C L S Q P I A S P N P L P L P I	TT 50 L W
G.R AKG	L A I V L P L	PAQ FLHS	CCTAAGICC TOOGITCA L S A W V H . V P G F I A K C L G S	P
N R A L I E L	E H . N T S	S L G S H W V	CCACGGITCT CTTCCATG T V L F H D P R F S S M ' H G S L P .	r
P W L L H G F	I E L . S Y	. H S N T H	CIGCAIGGIC CAAGATIC L H G P R F I C M V Q D S A W S K I P	H .
S L E P W N	S V R P P . D	RTP QEPQ	AGGICAGAGA ACACAAGG G Q R T Q G V R E H K A R S E N T R	L
P P C C H H V	W K Q G S S	P T T I	LEAARH FWKRPA FGSGPP	T
Y L G S	S G S A L G A	K D P R T P	CAGGIAACAA TTIGGIGA Q V T I W . R . Q F G D G N N L V T	P H
R R D E G T	L N P Q	P.R NHEG	GCATCICCAA AGCAATIG D L Q S N W I S K A I G G S P K Q L	K

25/33 -24-/-32 FIG 10 (continued) FIG 10B

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
AATGITCCIC	CCAAGGCAAA	AATGCCCCTA	AGATGRATIC	TGGAGAATTG	450
C S S	QGK	N A P K	MYS	GEL	
N V P P	K A K	M P L	R C I L	E N W	
M F L	P R Q K	CP.	D V F	WRIG	
GGACCAATCT	GACCCTCAGA	CAGTAAGAAA	AAAAATGACT	TATATICTIC	500
G P I .	P S D	s k k	K N D L	Y S S	
DQS	DPQT	V R K	K M T	YILL	
TNL	T L R	Q.EK	K . L	IFF	
TGCAGTACCG	CCTGGCCACG	GATATCCTCT	TCAAGGGGGA	GAAACCTGGC	550
AVP	P G H G	Y P L	QGG	ETWP	
QYR	LAT	DILF	KGE	KPG	
		I S S			
CTCCTGAGGG	AAGTATAAAT	TATAACACCA	TCTTACAGCT	AGACCIGITT	600
PEG	SIN	Y N T I	LQL	DLF	
LLRE	v . I	I T P	SYS.	T C F	
S.G	K Y K L	. н н	LTA	RPVL	
TGIAGAAAAG	GAGGCAAATG	GAGIGAAGIG	CCATATTIAC	AAACITICIT	650
CRKG	G K W	SEV	PYLQ	TFF	
VEK	E A N G	V K C	H I Y	K L S F	
		E . S A			
TTCATTAAAA	CACAACICCC	AATTATGTAA	ACAGIGIGAT	TIGIGICCIA	700
SLK	D N S Q	L C K	QCD	LCPT	
н. к	TTR	NYVN	s v i	CVL	
FIKR	QLA	I M .	T V . F	V S Y	
					•
CAGGAAGCCC	TCAGATCIAC	CICCCIACCC	CCCATCTCC	CIGACICCIT	750
G S P	QIY	L P T P	A S P	. L L	
QEAL	RST	S L P	R H L P	DSF	
R K P	S D L P	P Y P	G I S	LTPS	
CCCCAACTAA	TAAGGACCCA	CTTCAGCCCA	AACAGTOCAA	AAGGACATAG	800
PQLI	R T H	F S P	N S P K	G H	
P N	. G P T	S A Q	T V Q	K D I	
		L Q P K			

名し33 25 / 32 FIG 11 A

10	20	30	40	50	
1234567890 123	34567890 12	34567890	1234567890	1234567890	
GCATTGATA GC G I D S ' A L I A H H	ACCCATCA GAY	IGGCCAAA A K W P N	TCATTATTTA S L F T H Y L	CTGGACCAGG G P G L D Q A	50
CCTTTCAAA AC L F K T F S K L P F Q N	I K Q S	IGP .GP	VKH .SM	A K E I P K K	100
TAATCCCCTG CC. I P C L . S P A I N P L P	IA M LSP C	F L Q S F	E N K R R T K	E Q A N R P	150
ATTACCCAGG GG I T Q G I L P R G Y P G E	KTG N RLA '	. I r R F	L P T W Y P H	PNV GQMS	200
CAGGGATTIC AGG R D F S G I S A Q G F Q 1	IY.	S G Q L G R	I L S Y F H	L V G W W L G	250
GAGICTICT CC S L L L G V F S E E S S P	VGQ L.DR	K R P K D	KR. PRGN	.RH KGT	300
TAATGAAATA AT N N N E I I M K . F	S Q I W	T S G L P	P R I T P G L	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	350

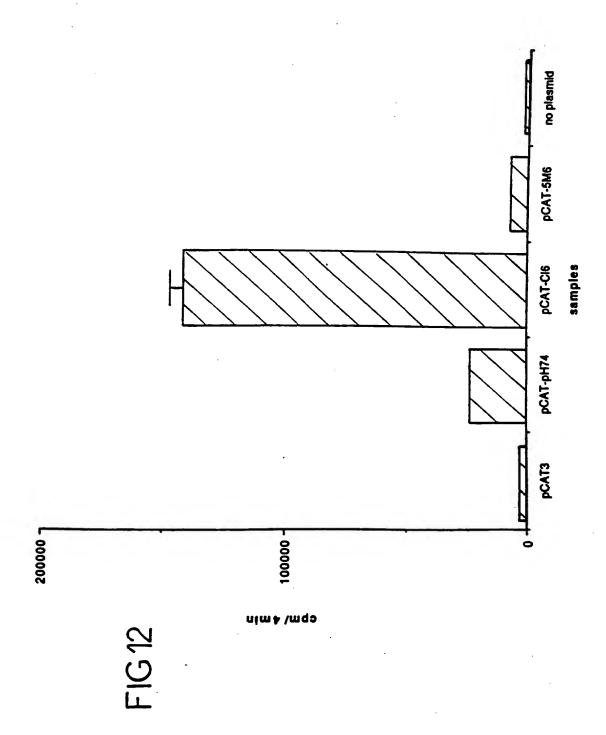


27|33 26 | 32 FIG 11 (continued) FIG 11B

	20 30 40 50	10
	1234567890 1234567890 1234567890 1234567890	1234567890
400	ITICAAGGCI GCAGIAACCC AGGGAGIAIC CCAGGIGITA	
	F Q G C S N P G S I P G V R	WPR
	F K A A V T Q G V S Q V L	G P A
	SRLQ.PREYPRC.	MAPL
450	ATCACTTACA CIGIGCCIGG AGGCCACAAT CCICCAGAAA	GGCATACAAT
450	S L T L C L E A T I L Q K	H T I
	HLH CAW RPQS SRK	GIOY
	TYT VPG GHN PPEK	AYN
	N I O O I N I P E K	
500	ATGAATGAAA CACTCAAAGA TCTAAAAAAG CTAACCCAAG	AGTCAAGAAA
	E.N TQR SKKA NPR	
	INET LKD LKK LTQE	
	. M K H S K I . K S . P K	S R K
	·	
550	GCATGACCT GITCIGITGC CLATAACCIT ACLAAGAATC	AAACCCACAT '
	MTCSVAYNL TKNP	NPH
	A.PVLLPITL LRI	T H I
	HDL FCC L . PY . ES	K P T L
		CD CD 1 CD 1
600	COCAAAAAG CAGGACTTAG CCCATACGAG ATGCTATATG	
	P K K Q D L A H T R C Y M	
	PKS RT. PIRD AIW	
	Q K A G L S P Y E M L Y G	T T 1
650	CIAACCAAT GACCTIGIGC TIGACTGAGA AATGGCCAAC	CATICACTURE (
050	. PM TLC LTEK WPT	
	LTN DLVL D.E MAN	WPF
700	CATCACCIC CITAGCCAAA TATCAACAAG TICITAAAAC	TIAGITICAG I
	SPP.PNINKFLKH	
	HHL LSQISTSS.N	
	I T S L A K Y Q Q V L K T	

28|33 27 / 32 FIG 11 (continued) FIG 11 C

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
ATCACAGGGA	ACCIGICCCC	GAGAGGAGGG	AAAGGAACTA	TICCACCCIG	750
H R E	PVP	ERRE	RNY	STL	_
ITGN	L S P	R G G	K G T I	PPW	
S Q G	T C P R	E E G	K E L	F H P G	
GIGACAIG					758
V T					
. н					
D M					



88

33

500

98

200

267

900

85

1100

88

500

534

1700 542 80

STITTEGAL TELAMANA AMMANA

500

467

TACCCTCAGG 9 7 8 9 הדהכוכודה אכוסהוכוכר האככככדה כפכוכדכאכך פכאככככרכ כאופכונוכים האכאככאכד אכרכככד 2 2 4 GIGCTICCTC TCATTCTTAG TGCCCCTAT OACCATCTAC Z () OUT CHENCE CONCRETE CONCRETE TREGITTE CHETTERARE GOODNETA GACAGIAC PACATO TCCTOACCT CATTICALE CATACCTCAA GOCAGACTAG **CCTGGTCAC** TITATITITA GACCTICAL TTGGACCCTG CATCACTAAG CARCOCCTAC COCA<u>TATAA A</u>CCAGGAT TCCAGCGC ACCACACE CCTTTGGT CCCTCCTT TGTATGGGC CTCTGTTTTC ACTCTATTC ACTCIATTAA CCANTCHAIG ACCTCACTAA AATGCTAATT AGGCAAAAT AGAGGTAAA GAAATAGCA ATCATCTATT GCCTGAGAGC ACAGCGGAA GGACAAGGAT 7 8 A . . ں 0 × + L C H H A H T H Y W T O K H I H F S C P TECATCHGET GOTTACKCC TECCACKCA ATACTCTCC CCCALACTOC ACCTICACTO CAGGTCHOG CAGAGAAA ACAGTAAG Q Q Q A R E K Q V K TACATGAAAC CCTCCGTACC CATACTCGCC OSTITICACE CANACETA CTANCISTIS SATOISCETE CECETOCACE TEAGECATA
V 8 A Q N P T N C M M C L P L H F R P Y GARCTETTO TITECAATET BOMATAACE CTCCTTITG TTATCAGAGE AGGAGTGCTA H + H TAAATGOTGA CATGGAACAG GTCACTGACT CTTOCTANCE GCEANAGAG GGGGAACCTO TCCKGAATTG TCACTGAGAA AGTTAAAGAA ATTGGAGATC GAATACAATG TAGAGCAGAG 8 Q t > CTCAGCCAAT GATGCCCTG GGTTCTCCCC TTCTTAGAAC CTCTAGCAGC TCTAATATTG TTACTCCTCT CICCITIGITA AGITTOTICIC TICCAGMIT GAGCTICIAA AGCTACAGA GITCITACAA ATIGAACCE AGATGAGTE COGCETACT ACCENTACT CEGATOTTA TOACATTOAA GOCACECETE CEGAGALAT CTCALCTOCA ACCTOCATT TOCTAGGODA ACCAMIGANTO COTAGGOTA GOTGOGANGO TOACTOCATO CACCOCOTA CATGOGGOTT GCAACTTAGO נוני # + 80 M X ST - - B OPLV SNL M R P O GTPPERI o u . . . CITTATCIA G ---0 2 ¥ ∀ √ CAGAATCTAT T B I H T T S V L V CTANGENCE ACANCHANG AGTACCATT CTOCANATAT TGATGCCCCA TCATATAGGA CATACCAGTA TOTCTGATGG GGOTGGAATT 2 8 7 GTACTGGCAT TGGCAGTATC ACAACCTCTA CTCAGTTCTA CTACAAACTA TCTCAAGAAA T G I G I T I S T G F Y Y K L S G K I ACTACTOCT CALANTCON CACCTITACA ر د د F L G P CTCTCMAAC 0 1 E H C L N G S S 8 A C ACCATACTA TAGACACCAC CACCTCCCAA CCCCTACAA AGGACTAGTT
P Y K G L V X > 0 S U & د د د o s Cap site 2 4 2 __ = S COCTCCATOA O s s 2 0 0 1 ANCTACTARA ACTICTACCAC TACTICACT * 4 + 0 CLACTICACC **CCTCAGCCTA** TCATGTGGTA CCCTACCACC GERRC TY V N THE AGCACCCTA S > r r 4 4 4 × × • . . ** ** > > = < --TOTALANTET V CACTIANCT 0 L N S S A G NTGCCCTCC CTTATCATAC ACCINGAGIT TCINTGAGA CCACCACTO TCTOTTCCAC CCCMCTAAC CODOCAAT 4 4 4 CCTOTTCCTO ACANTOGA ACTGACAG ATTTATACA r c < NTITANTACE ACCENCACTE **OTCTOTOOTA** ATCACCOTO DACCCTOCA **+** 0 0 A ACACCOANCO CTOCGOCCTC 2 2 0 --. . . . = 0 = ACCTOACCTO 1 # · > · PATOCCOCOC DATE 0 E I PATOCCOCAA 100 AATATTTTT CTTCCALCAT ** ** PATCTTAMC **M** 0 2 + = K L - -001

FIG18

400 700 800 900 1000 1100 170 1200 1300 237 1400 1500 1600 137 270 1700 20 203 303 337 370 1800 403 437 2000 470 TITATGGCTC AAGCTGAGCT TITGITGGCC AICCACCACT GCTGTTTGCC ACCGTCACAG ACCGGCTGCT GACTTCCATC CCTFTGGATC CAGCAGAGTO CCTTGGAATC CGTGAGACCA AGAACCCCAG GTCAGAGAAC ACAAGGCTTG CCACCATGTT GGAAGCAGCC CACCACCATT TTGGAAGCAG CCCGCCACTA TCTTGGGAGC TCTGGGAGCA AGGACCCCAG GTAACAATTT GGTGACCACG AAGGGACCTG AATCCGCAAC CATGAAGGGA TCTCCAAAGC AATGGGAAAC E F G ATCTTTGGTA TCTCAGTCAG GCCAACAATA GGATGACAAC AGAGGAAAGA ACAACTCCCA CAGGCCAGCA GGCAGTTCCC AGTGTAGACC CTCATTGGGA ATGTCCACTA TAACACAGGG AAAGGAAGAA AATCTTACTG CTTTTCTGGA CAGACTAAGG GAGGCATTGA GGAAGCATAC CTCCCTGTCA CCTGACTCTA CATGCAACTG | CACTCTTCTG GTCCGTGTTT AGATTCCATT GTTCCCCCCO AGGCAAAAT GCCCCTAGAA CGTATTCTGG AGAATTGGGA CCAATGTGAC ACTCAGACGC TAAGAAAGAA ACGATTTATA TTCTTCTGCA TCTTCTGTAG aaaggaggg aaatggagtg aagtgccata tgtgcaaact ttcttttcat taagagacaa ctcacaatta tgtaaaaagt gtggtttatg ccctacagga acaaaggggt aaacaatgaa ccaaagagtg ccaatattec ccgattaatgc ccctccaag cagtgaggg aggagaattc ggcccagcca gagtgcctgt TCCTTTGATC TGACATGGAG AGATATAATG TTACTACTAA ATCAGACACT AACCCCAAAT GAGAGAAGTG CCGCTGTAAC TGCAGCCCGA GAGTTTGGCG CACAGAATCA GAACATGGAG ATTGGTGCCA CAAACATTTG CTAACTTGCG TGCTAGAAGG ACTGAGGAAA ACTAGGAAGA AGCCTATGAA TTACTCAATG AGCCCTCAGA GTCCACCTCC CTACCCCAGC GTCCCTCCC CGACTCCTTC CTCAACTAAT AAGGACCCCC CTTTAACCCA AACGGTCCAA AAGGAGATAG KEID accitititict cictcagact taaagcaaat taaaatagac ctaggtaaat tctcagataa ccctgacggc tatattgatg titttacaagg gitaggacaa F C R V P V PDSI TTGAAGGCCA ACTAATCTTA AAGGATAAGT TTATCACTCA GTCAGCTGCA GACATTAGAA AAAAACTTCA AAAGTCCGTC TTAGGCTCGG AACAAAACTT Ω × H TIGAACTIGG CAACCICGGT TITITATAAT AGAGATCAGG AGGAGCAGGC AGAA#GGGAC AAATGGGATA AAAAAAAAAG GGCCACCGCT L N L A T S V F Y N R D Q E E Q A E W D K W D K K K R A T A CCCTCAGGCA AGCGGACTTT GGAGGCTCTG GAAAAGGGAA AAGCTGGGCA AATAGGAAGC CTAATAGGGC TTGCTTCCAG TGCGGTCTAC AAGGACACIT TAAAAAAGAT TGTCCAAATA GAAATAAGCC GCCCCTTGT CCATGCCCCT TACGTCAAGG GAATCACTGG AAGGCCCACT GCCCCAGGG P M N Y S M Y I D V L Q G L G Q OLDL SVDP G P A R TUCTIGATICEA GCACAGGGGG CCATTIGCCTC TUCCAATTIGG GUTAAAGGUT TGUCATTIGTT CUTGUACAGU TAAGTIGCUTG AACACTAGTC ACTGGGTTCC ACGGTTCTCT TCCATGACCC ATGGCTTCTA ATAGAGCTAT AACACTCACT GCATGGTCCA GTACCGCCTG GCCACATAT CCTCTTCAAG GGAGAGAAAC CTGGCTTCCT GAGGGAAGTA TAAATTATAA CATCATCTTA CAGCTAGACC S L S LGSE RFI A A R X X KAHC ر 1 1 T R K K C X X ERSAAVT K W D K LTQ ж ж N Y N I I L G 8 GQQAVP R L R E A L R K H T N R A K S V W.H.W TOTL CAGCAACCCC CTITGGGTCC CCTCCCAITG TATGGGAGCT CTGTTTTCAC TCTATTTAAAF PLQAVRG N R K K D P P SOL P D G LEGLRK E G S I TTPT N Q S R D N SAADIRK R N K P P P C/R P C P L STS T & 0 0 0 S W.A ITCAAGAIAC ICIGAGICAG AAGCCAITAA CCAGAIGAIC CAGCAGCAGG ACIGA FFSL KHL LTCV NLTAFLD O X N L G K F R L C T P S D T W L ख ख स K G K RILE VPSP PKSANIP DIMLLLN H T ж В о Н 0 8 0 K Q I K I D P L Q G LSQANNR E H G D W C H LILKDKP × a > ж ы л П Y P S A D P z a 네 K & S E M N N L S D. L X X P 0 4 ط ط ط X o di SFDL TCCGCTGTGC **AATCGAGCTG** r X V P P B SPOS M S T I AGAAACCCTA C V M A TAT > 0 × TAGTCATGG 0 0 0 S E S T B S E T

487

2055

3 7	200	700	134	500	200	234	267	900	1000	1100	1197
2 1 1 2	TAGGCATTGA G I D	ALGULATAAT R H H	AAATCACAGG K S Q G	AGCACTAGT A L V	CCAGGCGTTA Q A L	CTAACCCAGG L T Q R	TOCTOTATOG	L K T	TCCCTAGACA S L D T	TCANCCTO K P W	ACCOTOR
D X	ACCTTCCACT T F I L	V S Q	CCACATOCCA H H P	GAGGTAATAA E V I K	AGGGAGTATC G V S	TCTAAACAAG L N K	CCATACAGAA P Y R K	ATCACCAGT	TAATTCCCA M S P	CACACTCGAA H T R I	TCMGCMCA K 0 0
T S R	CCTTCAGCAA	ACCCCTCTC	TAGATTETAT	MAGTICCAA K F Q	ACAGTAACCC T V T Q	TCAAACGACA K R H	ACCACTENGE G L S	TENGCCAAAT L A K Y	AAAGAAATAT TOCACCTGG TGTCATGGTA TTAGTCAAGT COCTTCOCTC TAATTCCCCA TOCCTAGACA K & I P H P G V M V L V K S L P S M S P S L D T	TATCTATCC AACTGCGGTT AAAGTGGCTG GAGTGGAGTC TTGGATACAT CACACTCGAA TCAAACCCTG S I P I H I H I R I K P W	CCACAACGCT ACTATHICT TICAACCTCT ACACCATCTO TECCTCCTCT TCAAGCAACA ACCOTCAAD NA SYFFFFFFFFF B D L C L L F K Q Q P .
Al Management of the Control of the	TACTTAMAC L K T	GCAGATAGTC Q I V	CACTGGCAAC D W Q L	GTAGGACAGA R T E	F K A	ATCAMPACAC N K T L	CCCAAAGGC	CATCACCTCC I T S	TENGTCANGE L V K S	CAGTGCAGTC V E S	AGACCATCTC E D L
AGALGAAGAA E E E E	CLATOCALAT Q W K L	AACTATCAA T I K	CCCAACAAAAA PRE	GCCTTCCCCT	ATCOTCCTCC G P A	CCTTCACANA V E K	CULUCTOTIC Q N S P	TAGTTGCAGA V A D	TOTCATOOTA V M V	ANASTOSCTO R V A G	TTGAACCTCT
CCCAGTACTC	GGCAGCTAAC A A N	GCCTTTTCA G L F K	CAGGCAATTA G N Y	TTGGGCAGAG W A E	CAGAGTGACA Q S D N	CCTCAGGGAA S G K	ACTANGAATC L R I	ATOGCCAACT H A H L	TCCACCCTOG H P G	AACTGCGGTT T A V	ACCIATITICS S Y P P
GAMCCMGC	ractititisct L L L	TACTOCACCA T G P	GACAAAGAA E Q R T	CTTTCACTGG	CTGAGGCTTA G L	AGGCCACAAT R P Q S	CTATAGCCTT I A L	Tanccadada D Q B		TATCTATCCC S I P	
TCCCTCCGG	GAGGAAAAA	ATCATTATT S L F	rcmcaaa P s a	TCCCTAGATA	TCGGACTTCC	CTGCACCTAG	CCICIOTION A L L S	ACCTICTECT L L L	AGAGGAGGAA E E G	CCAGTCATTT P V I L	MANTECAGG
TATOCCCTAA N G N	AGCCACTGAA A T E	TAGCACCCAT CAGATAGCCA AATCATTATT TACTGGACCA GGCCTTTTCA AAACTATCAA GCAGATAGTC AGGGCCTGTG AAGTGTGCCA AAGAAATAAT S T $\overline{\mathbf{z}}$ Q I λ R λ $\underline{\underline{\mathbf{c}}}$ R λ G I λ R λ $\underline{\underline{\mathbf{c}}}$ R λ R λ R λ R λ λ R λ λ R λ λ R λ R λ R λ R R λ R R R R	CCCCTGCCTT ATCGCCAAGC TCCTTCAGGA GAACAAAGAA CAGGCAATTA CCCAAGAGAA GACTGGCAAC TAGATTTTAT CCACATGCCA AAATCACAGG ${f P}$ ${f P}$ ${f R}$ ${f Q}$ ${f R}$	GATTICAGTG TCTACTAJTC TGGGTAGATA CTTTCACTGG TTGGGCAGAG GCCTTCCCCT GTAGGACAGA AAAGTTCCAA GAGGTAATAA AGGCACTAGT F Q C L L 7 w V D T F T G w A E A F P C R T E K F Q E V I K A L V	TCATGAAGTA ATTCCCAGAT TCGGACTTCC CTGAGGCTTA CAGAGTGACA ATGGTCCTGC TTTCAAGGCC ACAGTAACCC AGGGAGTATC CCAGGGGTTA HEVER I	Gotatagnat atcacttaca ctgcacctag aggccacaat cctcagggaa gottgagaaa atga $\#$ cac tcanacgaca tctanacaag ctancceagg of \mathbf{z} or \mathbf{z} in \mathbf	AMCCCACCT COCATGOTCT GCTCTGTTGT CTATAGCCTT ACTAAGAATC CAUAACTCTC CCCAAAAGGC AGGACTTAGC CCATACAGAA TOCTGTATGG T H L A M S A L L S I A L L R I Q N S P Q K A G L S P Y R M L Y G	ACGGECTIC CEAACCAATG ACCTICTGCT TGACCAAGT TAGTTGCAAA CATCACCTCC TTAGCCAAAT ATCAACAAGT TCTTAAAACA R S P L T H D L L L D Q B K A H L V A D I T S L A K Y Q Q V L K T	TINCHAGGAG CETGTCCCCG AGAGGAGGAA	CATCCTOGGO AGGACCCTAC CCAGTCATTT S N G G P T P V I L	CATACTOCCG AAGGAACCCG AAAATCCAGG
gaaccegtag tatgagataa teceeteega gaaaccaage eceagtaete agaagaaaa atagaataga gaaceteaga agaacatagat thefeeeete $a_{\parallel}P$ v v $_{\parallel}W$ o $_{\parallel}P$ v v $_{\parallel}W$ o $_{\parallel}P$ v v $_{\parallel}W$ o $_{\parallel}P$ s $_{\parallel}P$ v v $_{\parallel}W$ o $_{\parallel}P$ s $_{\parallel}P$ v v $_{\parallel}W$ o $_{\parallel}P$ s $_{\parallel}P$ s $_{\parallel}P$ s $_{\parallel}P$ s	CAGGATGGCT AGCCACTGAA GAAGGAAAAA TACTTTTGCT GGCAGCTAAC CAATGGAAAT TACTTAAAAC CCTTCAGAA ACCTTCCACT TAGGCATTGA ${ m G}$ ${ m M}$ ${ m L}$	TAGCACCCAT S T H	cccracerr	SATTICAGIG	TCATGAAGTA H E V	CCTATAGAAT O I E Y	AACCCACCT T H L	ACGGCCTTC R S F	TTACAAGGAG L Q G A	CATCCTGGG	GATACTGCCG

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FIG 16